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Attorney Docket No. 005430.00002

PATENT

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Application of ) **BOX PCT**  
Matthias RÜSING, et al. )  
Serial No. Unassigned ) National Phase Application  
Filed: CONCURRENTLY HEREWITH ) PCT/EP00/08778  
 ) Filed: September 8, 2000  
 ) Group Art Unit: Unassigned  
For: NUCLEIC ACID WHICH IS OBTAINED )  
FROM TETRAHYMENA AND WHICH )  
CODES FOR A DELTA-6-DESATURASE, )  
THE PRODUCTION THEREOF AND USE )

SUBMISSION OF SEQUENCE LISTING

Assistant Commissioner of Patents & Trademarks  
Washington, D.C. 20231

Dear Sir:

A paper copy of a substitute sequence listing is submitted herewith to place the sequence listing consistent with U.S. practice. I believe that the sequences in the original application and the attached substitute sequence listing are identical. Substitution of the substitute sequence listing therefore adds no new matter to the specification

Applicants also submit a computer readable form of the formal sequence listing for use in the present application.

The content of the two forms, paper and CRF, are believed to be identical.

Respectfully submitted,

March 8, 2002

Date

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Boston, Massachusetts 02109  
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*Edm. Hummerding, Reg. No. 42653*  
for Peter D. McDermott  
Reg. No. 29,411

SEQUENCE LISTING

<110> Aventis Research & Technologies GmbH & Co KG

<120> Nucleic Acid Which is Obtained from Tetrahymena and which Codes  
delta 6-Desaturase, the Production Thereof and Use

<130> Banner & Witcoff Attorney Docket Number 005430.00002; National Phase  
Application of PCT/EP00/08778

<140> TBA

<141> 2002-03-08

<150> DE 19943270.8

<151> 1999-09-10

<160> 19

<170> PatentIn Ver. 2.1

<210> 1

<211> 1219

<212> DNA

<213> Tetrahymena thermophila

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aatatgactg cactgaatat gctaaatcaa ataagcatcc tggcggctct aatttcctca 180  
atttgtttat tgatgagaag taagatttga ctgaatatct cagaacactc cattctaagt 240  
aggctttgaa aattttaaaa tccttcctca agactggcgc aaaataagag gagactgaat 300  
ctcaaagag atttcaata ttaaagaaaa agcttaagca ttattcgaa ccaaactggc 360  
ctatcgaaat tggttattc ttaactacct ttactttatt tgcactgga tgttgactc 420  
aaaagtggta ttctctatt ccccttcttg tcttaatgca aatcatcagt ggttgattg 480  
gtcactctat gaaccacaat cgtaacccta tattaagaaa attcgcttta gtctacgctc 540  
ctctttgtgg tggttctct aataaatggt ggggtaggaa gcacaatcaa catcatatgt 600  
tcacaaacaa cattctaaag gacgaagata tctaacacga ttacaaattg tggttaattcc 660

ccttcttatt tttaaagtgg aaattagact ccatcttagc ttcttattat gaatttgaag 720  
gaatcttcct tgccttgac tgggtattat tattcaacta aaacttctat atcgtaattc 780  
tttctgaatt gattgctggt ttcttcagtg ctctattct tgttggaat catgaaaatg 840  
aaatgaaatt cgaaagaaga atcaccttac cattttcga acatcaaata gctgcaagca 900  
gaaactacgc ttccacgac atattctctc tacttattat gggtggtatg taatattaga 960  
ctgaacatca cttttccca taaattcctt tctacagatt acccaaagct cgtgtcataa 1020  
ttgctgaaga attaaagaag tggaacctta agattcatga aggacctatt ttgaaaaat 1080  
ctcaccttg aaaataaata aatttattt aaatgcatat ttattagta atactaacia 1140  
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<210> 2  
<211> 352  
<212> PRT  
<213> Tetrahymena thermophila

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1 5 10 15  
  
Glu Leu Leu Asn Glu Tyr Lys Phe Ile Tyr Lys Asp Thr Glu Tyr Asp  
20 25 30  
  
Cys Thr Glu Tyr Ala Lys Ser Asn Lys His Pro Gly Gly Leu Asn Phe  
35 40 45  
  
Leu Asn Leu Phe Ile Asp Glu Lys Gln Asp Leu Thr Glu Tyr Phe Arg  
50 55 60  
  
Thr Leu His Ser Lys Gln Ala Leu Lys Ile Leu Lys Ser Phe Pro Lys  
65 70 75 80  
  
Thr Gly Ala Lys Gln Glu Glu Thr Glu Ser Ser Lys Arg Phe Ser Ile  
85 90 95  
  
Leu Lys Lys Lys Leu Lys His Leu Phe Glu Pro Asn Trp Pro Ile Glu

100	105	110	
Ile Gly Leu Phe Leu Thr Thr Phe Thr Leu Phe Val Thr Gly Cys Leu			
115	120	125	
Thr Gln Lys Trp Tyr Phe Ser Ile Pro Leu Leu Val Leu Met Gln Ile			
130	135	140	
Ile Ser Gly Trp Ile Gly His Ser Met Asn His Asn Arg Asn Pro Ile			
145	150	155	160
Leu Arg Lys Phe Ala Leu Val Tyr Ala Pro Leu Cys Gly Gly Phe Ser			
165	170	175	
Asn Lys Trp Trp Gly Arg Lys His Asn Gln His His Met Phe Thr Asn			
180	185	190	
Asn Ile Leu Lys Asp Glu Asp Ile Gln His Asp Tyr Lys Leu Trp Gln			
195	200	205	
Phe Pro Phe Leu Phe Leu Lys Trp Lys Leu Asp Ser Ile Leu Ala Ser			
210	215	220	
Tyr Tyr Glu Phe Glu Gly Ile Phe Leu Ala Leu His Trp Val Leu Leu			
225	230	235	240
Phe Asn Gln Asn Phe Tyr Ile Val Ile Leu Ser Glu Leu Ile Ala Gly			
245	250	255	
Phe Phe Ser Ala Ser Ile Leu Val Gly Asn His Glu Asn Glu Met Lys			
260	265	270	
Phe Glu Arg Arg Ile Thr Leu Pro Phe Phe Glu His Gln Ile Ala Ala			
275	280	285	
Ser Arg Asn Tyr Ala Phe His Asp Ile Phe Ser Leu Leu Ile Met Gly			
290	295	300	

Gly Met Gln Tyr Gln Thr Glu His His Phe Phe Pro Gln Ile Pro Phe  
305                    310                    315                    320

Tyr Arg Leu Pro Lys Ala Arg Val Ile Ile Ala Glu Glu Leu Lys Lys  
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Trp Asn Leu Lys Ile His Glu Gly Pro Ile Phe Glu Lys Ser His Leu  
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<210> 3

<211> 2492

<212> DNA

<213> Tetrahymena thermophila

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ataattcgat tcgtgtaaga tggaaattga aagaattaag gtttagaaaa gttctttttg 180  
taaaataata gagttaaagt caataaattt tatattacgt aaatcttaaa gtgtgcaaat 240  
gttatcatta acaattctaa atgatgcaaa atatttaaat tattaataat aatgatagtt 300  
aataaaatca atatttcata ataataataa ggtatctatc tatctatcaa tattcaata 360  
aatattaatt aaaagggtat aaaataagta agcaaactaa atttaaaaaa caagcattat 420  
gggagttgat aagacttaag aagaaattgt tcttgaaaat aaacccgaac ttctcaacga 480  
atacaaattt atttacaagg atactgaata tgactgcact gaatatgcta aatcaaataa 540  
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atatttcaga acactccatt ctaagtaggc ttgaaaatt ttaaaatcct tcctaagac 660  
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taagcatgta aatacattca aatgatatct ttattgagca tatttagcat aattgataa 780  
tttcataag catattttaa attataaaaa tgaacatatt tttaaattaa tttagttatt 840  
cgaaccaaac tggcctatcg aaattgggtt attcttaact accttactt tatttgtcac 900  
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 agtcatatt tgatgctta atagtacaaa caatattga ttgtatgat taaattata 1200  
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 atacaaataa ttgaaaaag ctaaacttt ttctattaa aattaattac aaattgtaa 1500  
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 gaaactaagt tgatggtgt atttttaat ttcttaatt aattgtgaa taaacgatga 2400  
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 gaccacgct gccctatagt gagtcgtatt ac 2492

<210> 4

<211> 10

<212> PRT

<213> Tetrahymena thermophila

<400> 4

Trp Trp Lys Trp Asn His Asn Ala His His

1

5

10

<210> 5  
<211> 13  
<212> PRT  
<213> Tetrahymena thermophila

<400> 5  
Gly Gly Leu Gln Phe Gln Ile Glu His His Leu Phe Pro  
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<210> 6  
<211> 20  
<212> DNA  
<213> Artificial Sequence

<220>  
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<400> 6  
tggtggaart ggamncayaa 20

<210> 7  
<211> 20  
<212> DNA  
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cgdggraana rrtgrgttc 20

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gaccacgcgt atcgatgtcg actttttttt ttttttttv 40

<210> 9

<211> 28

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence:Primer

<400> 9

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<210> 10

<211> 29

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence:Primer

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<212> DNA

<213> Artificial Sequence



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<223> Description of Artificial Sequence:Primer

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29

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<213> Artificial Sequence

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<223> Description of Artificial Sequence:Primer

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<210> 13

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<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence:Primer

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<210> 14

<211> 27

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence:Primer

<400> 14

cttaagtctt atcaactccc ataatgc

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<210> 15

<211> 30

<212> DNA

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<223> Description of Artificial Sequence:Primer

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gaagtggaac cttaagattc atgaaggacc

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<210> 16

<211> 30

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence:Primer

<400> 16

gcattatgca tgttgataag acttaagaag

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<210> 17

<211> 35

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence:Primer

<400> 17

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<210> 18

<211> 30

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence:Primer

<400> 18

aaaaataaaa aagttgaaa aaaaaccttc

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<210> 19

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<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence:Primer

<400> 19

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22